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Pro Arg Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys Glu Val Lys Lys
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Gly His His Asn His Pro Lys Pro Gln Pro Asn Arg Arg Leu Ala Ala
     50                  55                  60

Gly Ala Val Pro Ser Ser Gln Ala Glu Glu Arg Tyr Asp Gly Val Ala
 65                  70                  75                  80

Pro Ile Glu Asp Lys Pro Ser Asn Ile Tyr Ser Asn Leu Cys Asn Gln
                 85                  90                  95

Ala His Ser Ala Gly Met Val Asp Asn Val Pro Gly Pro Ala Ser Asp
            100                 105                 110

Asp Asp Val Asp Ala Gly Gly Gly Arg Pro Xaa Pro Gly Gly Met Thr
        115                 120                 125

Xaa Met Met Met Met Thr Xaa Xaa Ser Lys Thr Gln Gly Lys Trp Asn
    130                 135                 140

Leu Pro Gly Xaa Xaa Cys Xaa Ala Leu Xaa Gly Val Asn Gln Ile Pro
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Glu Pro Xaa Xaa Pro Phe Xaa Lys Leu Phe
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<213> Zea mays

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Pro Pro Pro Phe Ile Thr Ser Phe Thr Glu Leu Leu Ser Gly Ala Gly
20 25 30

Asp Met Leu Gly Ala Gly Ala Asp Gln Glu Arg Ser Ser Pro Arg Gly
35 40 45

Leu Phe His Arg Gly Ala Arg Gly Val Pro Lys Phe Lys Ser Ala Gln
50 55 60

Pro Pro Ser Leu Pro Ile Ser Pro Pro Pro Met Ser Pro Ser Ser Tyr
65 70 75 80

Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser Pro
85 90 95

Val Leu Leu His Ser Ser Ser Asn Ile Leu Ala Ser Pro Thr Thr Gly
100 105 110

Ala Ile Pro Ala Gln Arg Phe Asp Trp Lys Lys Ala Ala Asp Leu Ile
115 120 125

Ala Ser Gln Ser Gln Gln Asp Gly Asp Ser Arg Ala Ala Ala Ala Gly
130 135 140

Phe Asp Asp Phe Ser Phe
145 150

<210> 5
<211> 1961

<212> DNA
<213> Oryza sativa

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<210> 6
<211> 488
<212> PRT
<213> Oryza sativa

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Pro Pro Phe Ile Thr Ser Phe Thr Glu Leu Leu Ser Gly Gly Gly Gly
20 25 30

Asp Leu Leu Gly Ala Gly Gly Glu Glu Arg Ser Pro Arg Gly Phe Ser
35 40 45

Arg Gly Gly Ala Arg Val Gly Gly Gly Val Pro Lys Phe Lys Ser Ala
50 55 60

Gln Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Val Ser Pro Ser Ser
65 70 75 80

Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Thr Glu Leu Leu Asp Ser
85 90 95

Pro Val Leu Leu Ser Ser Ser His Ile Leu Ala Phe Pro Thr Thr Gly
100 105 110

Ala Ile Pro Ala Gln Arg Tyr Asp Trp Lys Ala Ser Ala Asp Leu Ile
115 120 125

Ala Ser Gln Gln Asp Asp Ser Arg Gly Asp Phe Ser Phe His Thr Asn
130 135 140

Ser Asp Ala Met Ala Ala Gln Pro Ala Ser Phe Pro Ser Phe Lys Glu
145 150 155 160

Gln Glu Gln Gln Val Val Glu Ser Ser Lys Asn Gly Ala Ala Ala Ala
165 170 175

Ser Ser Asn Lys Ser Gly Gly Gly Gly Asn Asn Lys Leu Glu Asp Gly
180 185 190

Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn
195 200 205

Pro Arg Ser Tyr Tyr Lys Cys Thr Tyr Asn Gly Cys Ser Met Lys Lys
210 215 220

Lys Val Glu Arg Ser Leu Ala Asp Gly Arg Ile Thr Gln Ile Val Tyr
225 230 235 240

Lys Gly Ala His Asn His Pro Lys Pro Leu Ser Thr Ala Ala Thr Pro
245 250 255

Leu Pro Ala Pro Pro Pro Pro Pro Ala Pro Thr Thr Ser Arg Arg Pro
260 265 270

Ala Arg Ala Arg Thr Ser Thr Pro Pro Arg Arg Pro Arg Thr Pro Pro
275 280 285

Ser Arg Ser Ala Thr Thr Arg Pro Thr Thr His Arg Thr Ala Ala Arg
290 295 300

Ala Thr Ser Pro Lys Pro Ser Ala Gly Lys Glu Asp Ala Asp Asn Glu
305 310 315 320

Gly Ser Ser Gly Gly Met Gly Gly Gly Ala Gly Gly Asn Pro Val Arg
325 330 335

Glu Pro Arg Leu Val Val Gln Thr Leu Ser Asp Ile Asp Ile Leu Asp
340 345 350

Asn Gly Phe Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys Gly Asn
355 360 365

Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Thr Val Gly Cys Pro Val
370 375 380

Arg Lys His Val Glu Arg Ala Ser His Asp Thr Arg Ala Val Ile Thr
385 390 395 400

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Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Val Arg Pro Arg Arg
                405                 410                 415

Arg Arg Arg Thr Arg Pro Gly Ala Gly Val Ala Tyr Gly Trp Gly Arg
            420                 425                 430

Ser Gly Pro Thr Asp Val Ala Ala Ala Gln Gln Gly Pro Tyr Thr Leu
        435                 440                 445

Glu Met Leu Pro Asn Pro Ala Gly Leu Tyr Gly Gly Tyr Gly Ala Gly
    450                 455                 460

Ala Gly Gly Ala Ala Phe Pro Arg Thr Lys Asp Glu Arg Arg Asp Asp
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Leu Phe Val Glu Ser Leu Leu Cys
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<210> 7
<211> 2086
<212> DNA
<213> Oryza sativa

<400> 7

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<210> 8
<211> 487
<212> PRT
<213> Oryza sativa

<400> 8

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             20                  25                  30

Asp Leu Leu Gly Ala Gly Gly Glu Glu Arg Ser Pro Arg Gly Phe Ser
         35                  40                  45

Arg Gly Gly Ala Arg Val Gly Gly Gly Val Pro Lys Phe Lys Ser Ala
     50                  55                  60

Gln Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Val Ser Pro Ser Ser
 65                  70                  75                  80

Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Thr Glu Leu Leu Asp Ser
                 85                  90                  95

Pro Val Leu Leu Ser Ser Ser His Ile Leu Ala Ser Pro Thr Thr Gly
            100                 105                 110

Ala Ile Pro Ala Gln Arg Tyr Asp Trp Lys Ala Ser Ala Asp Leu Ile
        115                 120                 125

Ala Ser Gln Gln Asp Asp Ser Arg Gly Asp Phe Ser Phe His Thr Asn
    130                 135                 140

Ser Asp Ala Met Ala Ala Gln Pro Ala Ser Phe Pro Ser Phe Lys Glu
145                 150                 155                 160

Gln Glu Gln Gln Val Val Glu Ser Ser Lys Asn Gly Ala Ala Ala Ala
                165                 170                 175

Ser Ser Asn Lys Ser Gly Gly Gly Gly Asn Asn Lys Leu Glu Asp Gly
            180                 185                 190

Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn
        195                 200                 205

Pro Arg Ser Tyr Tyr Lys Cys Thr Tyr Asn Gly Cys Ser Met Lys Lys
    210                 215                 220

Lys Val Glu Arg Ser Leu Ala Asp Gly Arg Ile Thr Gln Ile Val Tyr
225                 230                 235                 240

Lys Gly Ala His Asn His Pro Lys Pro Leu Ser Thr Arg Arg Asn Ala
                245                 250                 255

Ser Ser Cys Ala Thr Ala Ala Ala Cys Ala Asp Asp Leu Ala Ala Pro
            260                 265                 270
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Gly Ala Gly Ala Asp Gln Tyr Ser Ala Ala Thr Pro Glu Asn Ser Ser
        275                 280                 285

Val Thr Phe Gly Asp Asp Glu Ala Asp Asn Ala Ser His Arg Ser Glu
    290                 295                 300

Gly Asp Glu Pro Glu Ala Lys Arg Trp Lys Glu Asp Ala Asp Asn Glu
305                 310                 315                 320

Gly Ser Ser Gly Gly Met Gly Gly Gly Ala Gly Gly Lys Pro Val Arg
                325                 330                 335

Glu Pro Arg Leu Val Val Gln Thr Leu Ser Asp Ile Asp Ile Leu Asp
            340                 345                 350

Asp Gly Phe Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys Gly Asn
        355                 360                 365

Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Thr Val Gly Cys Pro Val
    370                 375                 380

Arg Lys His Val Glu Arg Ala Ser His Asp Thr Arg Ala Val Ile Thr
385                 390                 395                 400

Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Val Gly Arg Gly Gly
                405                 410                 415

Gly Gly Gly Arg Ala Pro Ala Pro Ala Pro Pro Thr Ser Gly Ala Ile
            420                 425                 430

Arg Pro Ser Ala Val Ala Ala Ala Gln Gln Gly Pro Tyr Thr Leu Glu
        435                 440                 445

Met Leu Pro Asn Pro Ala Gly Leu Tyr Gly Gly Tyr Gly Ala Gly Ala
    450                 455                 460

Gly Gly Ala Ala Phe Pro Arg Thr Lys Asp Glu Arg Arg Asp Asp Leu
465                 470                 475                 480

Phe Val Glu Ser Leu Leu Cys
                485
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<210> 9
<211> 1928
<212> DNA
<213> Glycine max

<400> 9

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ctccttcttc ttactttgct attcctcctg gtttgagccc tgctgagctt cttgactcgc  300
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cataccagtc ttcaaatgtc acagttcaaa cacaacagcc atggagtttt caggaggcca  540
cgaaacagga taatttttcc tcaggaaagg gtatgatgaa aactgaaaac tcttcttcca  600
tgcagagttt ttcccctgag attgctagtg tccaaactaa ccatagcaat gggtttcaat  660
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<210> 10
<211> 575
<212> PRT
<213> Glycine max

<400> 10

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Phe Thr Asn Phe Thr Phe Ser Thr His Pro Phe Met Thr Thr Ser Phe
             20                  25                  30

Ser Asp Leu Leu Ala Ser Pro Leu Asp Asn Asn Lys Pro Pro Gln Gly
         35                  40                  45

Gly Leu Ser Glu Arg Thr Gly Ser Gly Val Pro Lys Phe Lys Ser Thr
     50                  55                  60

Pro Pro Pro Ser Leu Pro Leu Ser Pro Pro Pro Ile Ser Pro Ser Ser
 65                  70                  75                  80

Tyr Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser
                 85                  90                  95

Pro Val Leu Leu Asn Ser Ser Asn Ile Leu Pro Ser Pro Thr Thr Gly
            100                 105                 110

Ala Phe Val Ala Gln Ser Phe Asn Trp Lys Ser Ser Ser Gly Gly Asn
        115                 120                 125

Gln Gln Ile Val Lys Glu Glu Asp Lys Ser Phe Ser Asn Phe Ser Phe
    130                 135                 140

Gln Thr Arg Ser Gly Pro Pro Ala Ser Ser Thr Ala Thr Tyr Gln Ser
145                 150                 155                 160
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Ser Asn Val Thr Val Gln Thr Gln Gln Pro Trp Ser Phe Gln Glu Ala
                165                 170                 175

Thr Lys Gln Asp Asn Phe Ser Ser Gly Lys Gly Met Met Lys Thr Glu
            180                 185                 190

Asn Ser Ser Ser Met Gln Ser Phe Ser Pro Glu Ile Ala Ser Val Gln
        195                 200                 205

Thr Asn His Ser Asn Gly Phe Gln Ser Asp Tyr Gly Asn Tyr Pro Pro
    210                 215                 220

Gln Ser Gln Thr Leu Ser Arg Arg Ser Asp Asp Gly Tyr Asn Trp Arg
225                 230                 235                 240

Lys Tyr Gly Gln Lys Gln Val Lys Gly Ser Glu Asn Pro Arg Ser Tyr
                245                 250                 255

Tyr Lys Cys Thr Tyr Pro Asn Cys Pro Thr Lys Lys Lys Val Glu Arg
            260                 265                 270

Ser Leu Asp Gly Gln Ile Thr Glu Ile Val Tyr Lys Gly Thr His Asn
        275                 280                 285

His Pro Lys Pro Gln Asn Thr Arg Arg Asn Ser Ser Asn Ser Ser Ser
    290                 295                 300

Leu Ala Ile Pro His Ser Asn Ser Ile Arg Thr Glu Ile Pro Asp Gln
305                 310                 315                 320

Ser Tyr Ala Thr His Gly Ser Gly Gln Met Asp Ser Ala Ala Thr Pro
                325                 330                 335

Glu Asn Ser Ser Ile Ser Ile Gly Asp Asp Asp Phe Glu Gln Ser Ser
            340                 345                 350

Gln Lys Cys Lys Ser Gly Gly Asp Glu Tyr Asp Glu Asp Glu Pro Asp
        355                 360                 365

Ala Lys Arg Trp Lys Ile Glu Gly Glu Asn Glu Gly Met Ser Ala Pro
    370                 375                 380

Gly Ser Arg Thr Val Arg Glu Pro Arg Val Val Val Gln Thr Thr Ser
385                 390                 395                 400

Asp Ile Asp Ile Leu Asp Asp Gly Tyr Arg Trp Arg Lys Tyr Gly Gln
                405                 410                 415

Lys Val Val Lys Gly Asn Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr
            420                 425                 430

His Pro Gly Cys Pro Val Arg Lys His Val Glu Arg Ala Ser His Asp
        435                 440                 445

Leu Arg Ala Val Ile Thr Thr Tyr Glu Gly Lys His Asn His Asp Val
    450                 455                 460

Pro Ala Ala Arg Gly Ser Gly Ser His Ser Val Asn Arg Pro Met Pro
465                 470                 475                 480
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Asn Asn Ala Ser Asn His Thr Asn Thr Ala Ala Thr Ser Val Arg Leu
485 490 495

Leu Pro Val Ile His Gln Ser Asp Asn Ser Leu Gln Asn Gln Arg Ser
500 505 510

Gln Ala Pro Pro Glu Gly Gln Ser Pro Phe Thr Leu Glu Met Leu Gln
515 520 525

Ser Pro Gly Ser Phe Gly Phe Ser Gly Phe Gly Asn Pro Met Gln Ser
530 535 540

Tyr Val Asn Gln Gln Gln Leu Ser Asp Asn Val Phe Ser Ser Arg Thr
545 550 555 560

Lys Glu Glu Pro Arg Asp Asp Met Phe Leu Glu Ser Leu Leu Cys
565 570 575

<210> 11
<211> 2158
<212> DNA
<213> Triticum aestivum

<400> 11
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ccggcgcggg cgcgaggtac aaggccatgt ccccggcgcg gctgccgatc tcgcgcgagc 180
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<210> 12
<211> 619
<212> PRT
<213> Triticum aestivum

<400> 12

Thr	Ser	Arg	Thr	Ala	Pro	Met	Ala	Asp	Ser	Pro	Asn	Pro	Ser	Ser	Gly
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Asp	Leu	Pro	Ser	Ala	Ala	Gly	Ser	Ser	Pro	Glu	Lys	Pro	Tyr	Pro	Ala
			20					25				30			
Asp	Arg	Arg	Val	Ala	Ala	Leu	Ala	Gly	Ala	Gly	Ala	Arg	Tyr	Lys	Ala
		35					40					45			
Met	Ser	Pro	Ala	Arg	Leu	Pro	Ile	Ser	Arg	Glu	Pro	Cys	Leu	Thr	Ile
	50					55					60				
Pro	Ala	Gly	Phe	Ser	Pro	Ser	Ala	Leu	Leu	Asp	Ser	Pro	Val	Leu	Leu
65					70					75					80
Thr	Asn	Phe	Lys	Val	Glu	Pro	Ser	Pro	Thr	Thr	Gly	Ser	Leu	Ser	Met
				85					90					95	
Ala	Ala	Ile	Met	His	Lys	Ser	Ala	His	Pro	Asp	Ile	Leu	Pro	Ser	Pro
			100					105					110		
Arg	Asp	Lys	Ser	Ile	Arg	Ala	His	Glu	Asp	Gly	Gly	Ser	Arg	Asp	Phe
		115					120					125			
Glu	Phe	Lys	Pro	His	Leu	Asn	Ser	Ser	Ser	Gln	Ser	Leu	Ala	Pro	Ala
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Met	Ser	Asp	Leu	Lys	Lys	His	Glu	His	Ser	Met	Gln	Asn	Gln	Ser	Met
145					150					155					160
Asn	Pro	Ser	Ser	Ser	Ser	Ser	Asn	Met	Val	Asn	Glu	Asn	Arg	Pro	Pro
				165					170					175	
Cys	Ser	Arg	Glu	Ser	Ser	Leu	Thr	Val	Asn	Val	Ser	Ala	Pro	Asn	Gln
			180					185					190		
Pro	Val	Gly	Met	Val	Gly	Leu	Thr	Asp	Asn	Met	Pro	Ala	Glu	Val	Gly
		195					200					205			
Thr	Ser	Glu	Pro	Gln	Gln	Met	Asn	Ser	Ser	Asp	Asn	Ala	Met	Gln	Glu
	210					215					220				
Pro	Gln	Ser	Glu	Asn	Val	Ala	Asp	Lys	Ser	Ala	Asp	Asp	Gly	Tyr	Asn
225					230					235					240
Trp	Arg	Lys	Tyr	Gly	Gln	Lys	His	Val	Lys	Gly	Ser	Glu	Asn	Pro	Arg
				245					250					255	
Ser	Tyr	Tyr	Lys	Cys	Thr	His	Pro	Asn	Cys	Glu	Val	Lys	Lys	Leu	Leu
			260					265					270		

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Glu Arg Ala Val Asp Gly Leu Ile Thr Glu Val Val Tyr Lys Gly Arg
        275                 280                 285

His Asn His Pro Lys Pro Gln Pro Asn Arg Arg Leu Ala Gly Gly Ala
    290                 295                 300

Val Pro Ser Asn Gln Gly Glu Glu Arg Tyr Asp Gly Ala Ala Ala Ala
305                 310                 315                 320

Asp Asp Lys Ser Ser Asn Ala Leu Ser Asn Leu Ala Asn Pro Val Asn
                325                 330                 335

Ser Pro Gly Met Val Glu Pro Val Pro Val Ser Val Ser Asp Asp Asp
            340                 345                 350

Ile Asp Ala Gly Gly Gly Arg Pro Tyr Pro Gly Asp Asp Ala Thr Glu
        355                 360                 365

Glu Asp Leu Glu Ser Lys Arg Arg Lys Met Glu Ser Ala Gly Ile Asp
    370                 375                 380

Ala Ala Leu Met Gly Lys Pro Asn Arg Glu Pro Arg Val Val Val Gln
385                 390                 395                 400

Thr Val Ser Glu Val Asp Ile Leu Asp Asp Gly Tyr Arg Trp Arg Lys
                405                 410                 415

Tyr Gly Gln Lys Val Val Lys Gly Asn Pro Asn Pro Arg Ser Tyr Tyr
            420                 425                 430

Lys Cys Thr Ser Thr Gly Cys Pro Val Arg Lys His Val Glu Arg Ala
        435                 440                 445

Ser His Asp Pro Lys Ser Val Ile Thr Thr Tyr Glu Gly Lys His Asn
    450                 455                 460

His Glu Val Pro Ala Ala Arg Asn Ala Thr His Glu Met Ser Ala Pro
465                 470                 475                 480

Pro Met Lys Asn Val Val His Gln Ile Asn Ser Ser Met Pro Ser Ser
                485                 490                 495

Ile Gly Gly Met Met Arg Ala Cys Glu Ala Arg Asn Phe Ser Asn Gln
            500                 505                 510

Tyr Ser Gln Ala Ala Glu Thr Asp Asn Val Ser Leu Asp Leu Gly Val
        515                 520                 525

Gly Ile Ser Pro Asn His Ser Asp Ala Thr Asn Gln Met Gln Ser Ser
    530                 535                 540

Gly Pro Asp Gln Met Gln Tyr Gln Met Gln Ser Met Ala Ser Met Tyr
545                 550                 555                 560

Gly Asn Met Arg His Pro Ser Ser Met Ala Val Pro Thr Val Gln Gly
                565                 570                 575

Asn Ser Ala Gly Arg Met Tyr Gly Ser Arg Glu Glu Lys Gly Asn Glu
            580                 585                 590
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Gly Phe Thr Phe Arg Ala Thr Pro Met Asp His Ser Ala Asn Leu Cys
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Tyr Ser Gly Ala Gly Asn Leu Val Met Gly Pro
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<212> PRT
<213> Ipomoea batatas

<400> 13

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Met Ala Ala Ser Ser Gly Thr Ile Asp Ala Pro Thr Ala Ser Ser Ser
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Phe Ser Phe Ser Thr Ala Ser Ser Phe Met Ser Ser Phe Thr Asp Leu
             20                  25                  30

Leu Ala Ser Asp Ala Tyr Ser Gly Gly Ser Val Ser Arg Gly Leu Gly
         35                  40                  45

Asp Arg Ile Ala Glu Arg Thr Gly Ser Gly Val Pro Lys Phe Lys Ser
     50                  55                  60

Leu Pro Pro Pro Ser Leu Pro Leu Ser Ser Pro Ala Val Ser Pro Ser
 65                  70                  75                  80

Ser Tyr Phe Ala Phe Pro Pro Gly Leu Ser Pro Ser Glu Leu Leu Asp
                 85                  90                  95

Ser Pro Val Leu Leu Ser Ser Ser Asn Ile Leu Pro Ser Pro Thr Thr
            100                 105                 110

Gly Thr Phe Pro Ala Gln Thr Phe Asn Trp Lys Asn Asp Ser Asn Ala
        115                 120                 125

Ser Gln Glu Asp Val Lys Gln Glu Glu Lys Gly Tyr Pro Asp Phe Ser
    130                 135                 140

Phe Gln Thr Asn Ser Ala Ser Met Thr Leu Asn Tyr Glu Asp Ser Lys
145                 150                 155                 160

Arg Lys Asp Glu Leu Asn Ser Leu Gln Ser Leu Pro Pro Val Thr Thr
                165                 170                 175

Ser Thr Gln Met Ser Ser Gln Asn Asn Gly Gly Ser Tyr Ser Glu Tyr
            180                 185                 190

Asn Asn Gln Cys Cys Pro Pro Ser Gln Thr Leu Arg Glu Gln Arg Arg
        195                 200                 205

Ser Asp Asp Gly Tyr Asn Trp Arg Lys Tyr Gly Gln Lys Gln Val Lys
    210                 215                 220

Gly Ser Glu Asn Pro Arg Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys
225                 230                 235                 240
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Pro Thr Lys Lys Lys Val Glu Arg Ala Leu Asp Gly Gln Ile Thr Glu
245 250 255

Ile Val Tyr Lys Gly Ala His Asn His Pro Lys Pro Gln Ser Thr Arg
260 265 270

Arg Ser Ser Ser Ser Thr Ala Ser Ser Ala Ser Thr Leu Ala Ala Gln
275 280 285

Ser Tyr Asn Ala Pro Ala Ser Asp Val Pro Asp Gln Ser Tyr Trp Ser
290 295 300

Asn Gly Asn Gly Gln Met Asp Ser Val Ala Thr Pro Glu Asn Ser Ser
305 310 315 320

Ile Ser Val Gly Asp Asp Glu Phe Glu Gln Ser Ser Gln Lys Arg Glu
325 330 335

Ser Gly Gly Asp Glu Phe Asp Glu Asp Glu Pro Asp Ala Lys Arg Trp
340 345 350

Lys Val Glu Asn Glu Ser Glu Gly Val Ser Ala Gln Gly Ser Arg Thr
355 360 365

Val Arg Glu Pro Arg Val Val Val Gln Thr Thr Ser Asp Ile Asp Ile
370 375 380

Leu Asp Asp Gly Tyr Arg Trp Arg Lys Tyr Gly Gln Lys Val Val Lys
385 390 395 400

Gly Asn Pro Asn Pro Arg Ser Tyr Tyr Lys Cys Thr Ser Gln Gly Cys
405 410 415

Pro Val Arg Lys His Val Glu Arg Ala Ser His Asp Ile Arg Ser Val
420 425 430

Ile Thr Thr Tyr Glu Gly Lys His Asn His Asp Val Pro Ala Ala Arg
435 440 445

Gly Ser Gly Ser His Gly Leu Asn Arg Gly Ala Asn Pro Asn Asn Asn
450 455 460

Ala Ala Met Ala Met Ala Ile Arg Pro Ser Thr Met Ser Leu Gln Ser
465 470 475 480

Asn Tyr Pro Ile Pro Ile Pro Ser Thr Arg Pro Met Gln Gln Gly Glu
485 490 495

Gly Gln Ala Pro Tyr Glu Met Leu Gln Gly Ser Gly Gly Phe Gly Tyr
500 505 510

Ser Gly Phe Gly Asn Pro Met Asn Ala Tyr Ala Asn Gln Ile Gln Asp
515 520 525

Asn Ala Phe Ser Arg Ala Lys Glu Glu Pro Arg Asp Asp Leu Phe Leu
530 535 540

Asp Thr Leu Leu Ala
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<210> 14
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Synthetic oligonucleotide

<400> 14
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